

Pat17 atggcaactactaaatcttttttaattttaatatattatgatattagcaactactagttca 60  
M A T T K S F L I L I F M I L A T T S S 20

Pat17 acatttgctcagttgggagaaatggtgactgttcttagtattgatggaggtggaattaga 120  
T F A Q L G E M V T V L S I D G G G I R 40

Pat17 gggatcattccggctaccattctcgaatttcttgaaggacaacttcaggaaatggacaat 180  
G I I P A T I L E F L E G Q L Q E M D N 60

Pat17 aatgcagatgcaagacttgcagattactttgatgtaattggaggaacaagtacaggaggt 240  
N A D A R L A D Y F D V I G G T S T G G 80

Pat17 ttattgactgctatgataagtactccaaatgaaaacaatcgaccctttgctgctgccaaa 300  
L L T A M I S T P N E N N R P F A A A K 100

Pat17 gaaattgtacctttttacttcgaacatggccctcagatttttaatcctagtggtcaaatt 360  
E I V P F Y F E H G P Q I F N P S G Q I 120

Pat17 ttaggcccaaaatatgatggaaaatatcttatgcaagttcttcaagaaaaacttggagaa 420  
L G P K Y D G K Y L M Q V L Q E K L G E 140

Pat17 actcgtgtgcatcaagctttgacagaagttgtcatctcaagctttgacatcaaaacaaat 480  
T R V H Q A L T E V V I S S F D I K T N 160

Pat17 aagccagtaatatcactaagtcaaatttagcaaactctccagaattggatgctaagatg 540  
K P V I F T K S N L A N S P E L D A K M 180

Pat17 tatgacataagttattccacagcagcagctccaacatattttcctccgcattactttgtt 600  
Y D I S Y S T A A A P T Y F P P H Y F V 200

Pat17 actaatactagtaatggagatgaatatgagttcaatcttgttgatgggtgctgttgctact 660  
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G T T S E F D K T Y T A K E A A T W T A 280

Pat17 gtacattggatggttagttatacagaaaatgactgatgcagcaagttcttacatgactgat 900  
V H W M L V I Q K M T D A A S S Y M T D 300

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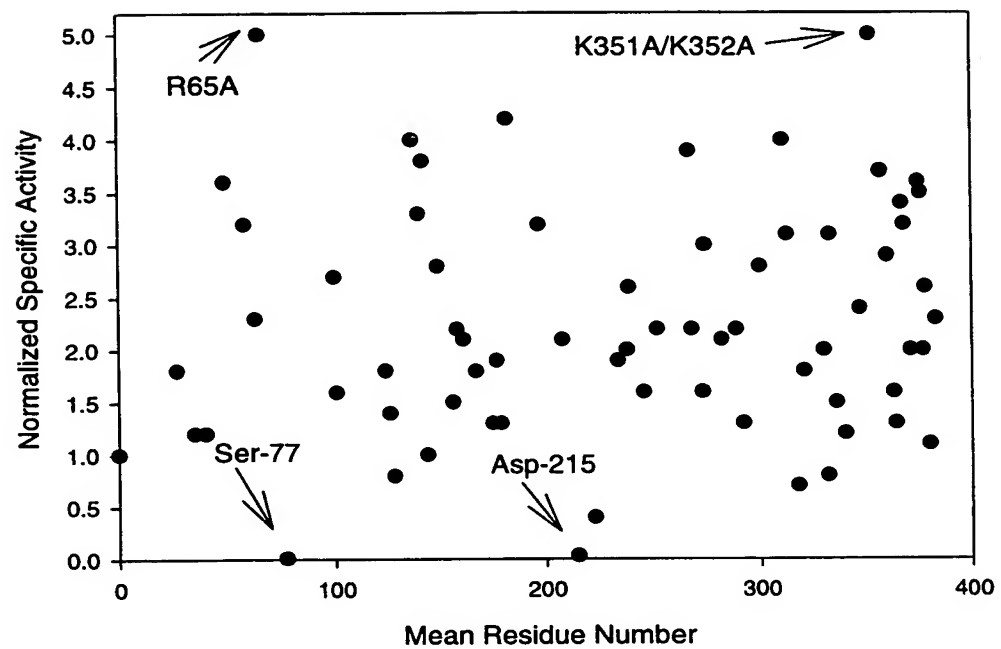
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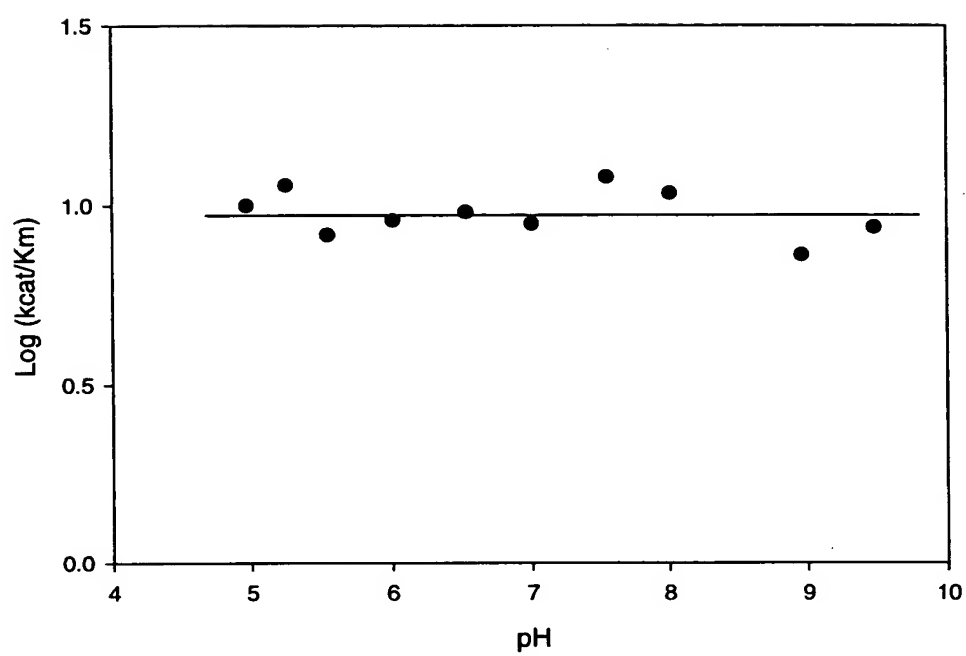
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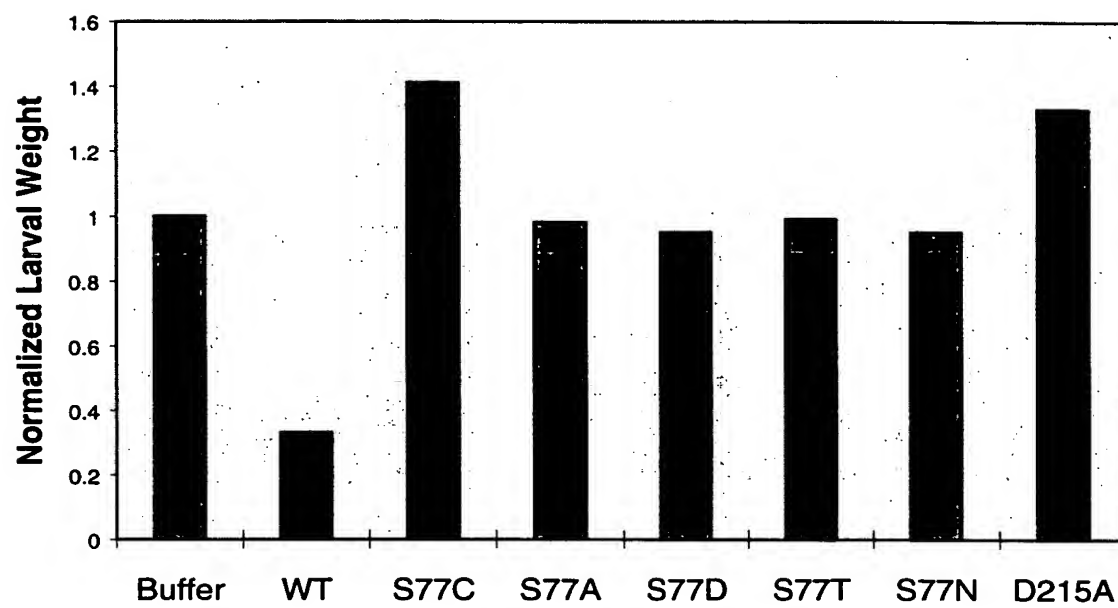
**Figure 1**



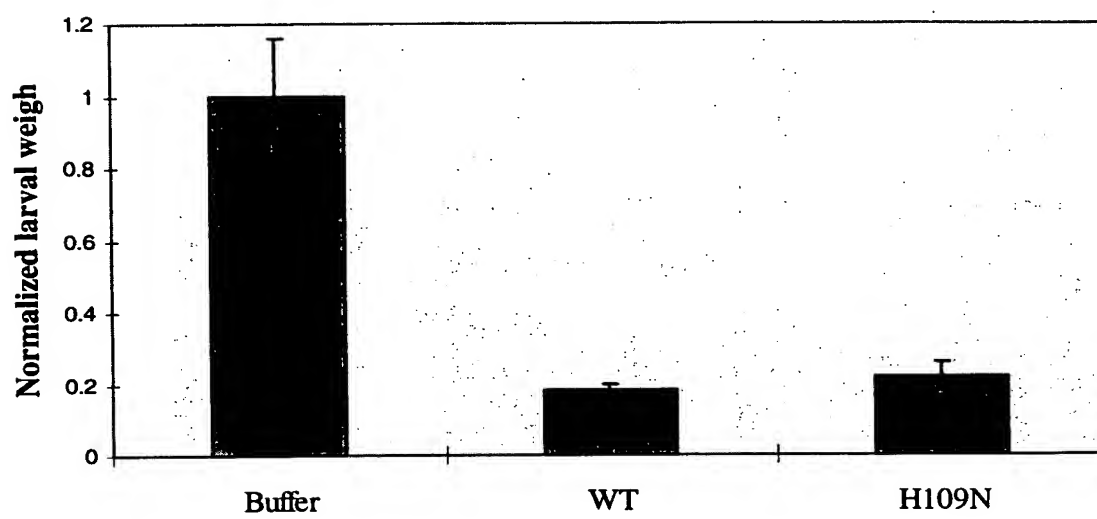
**Figure 2**



**Figure 3**



**Figure 4**



**Figure 5**



**Figure 6a**

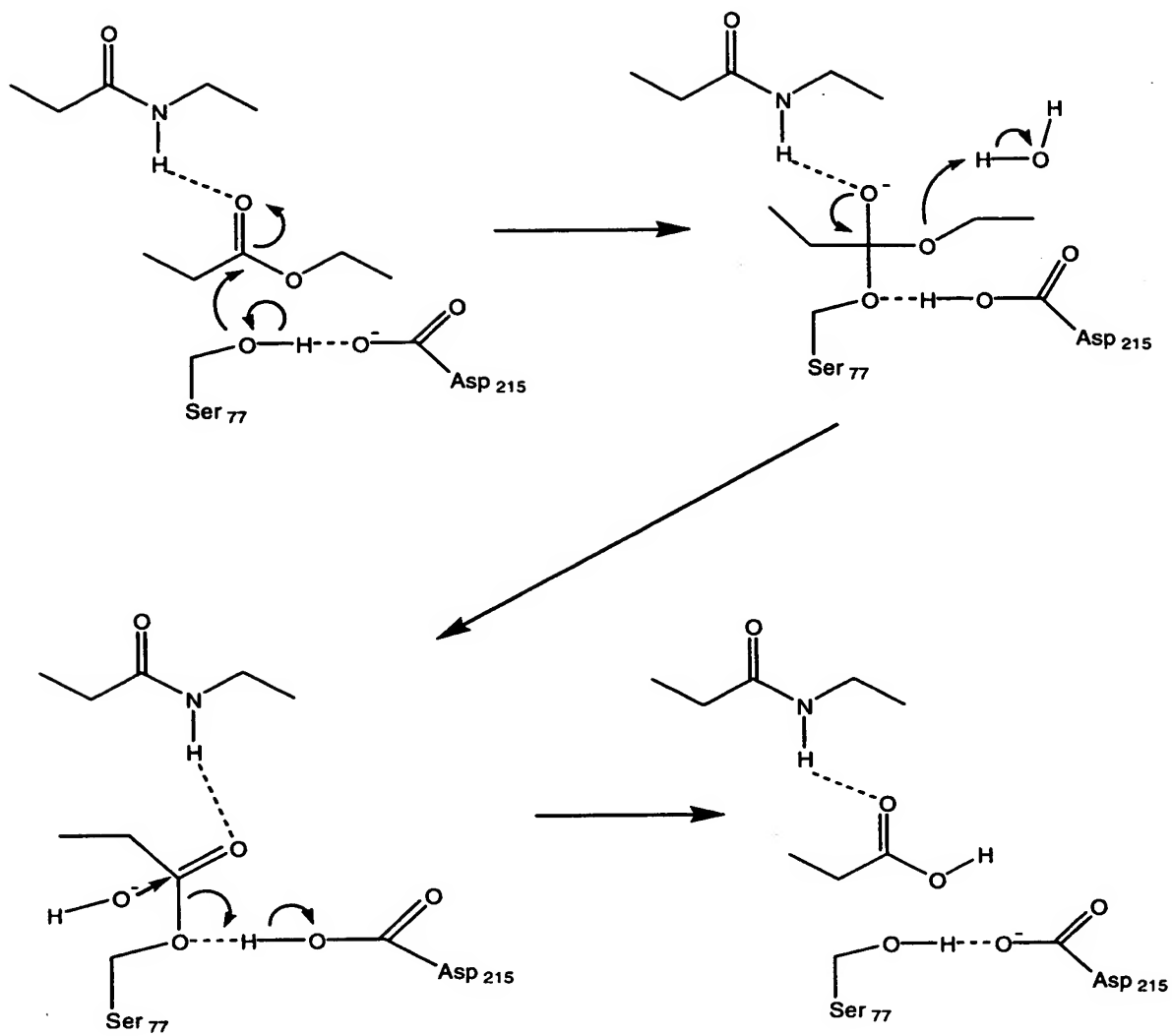


**Figure 6b**



**Figure 7**





**Figure 8**

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gi|PatFm -----MALEEM 6
gi|PatIm -----PWLEEM 6
gi|PatL+ -----MATTKSFLILFFMILA-----TTSSTCAKLEEM 28
gi|PatA+ -----MATTKSFLILFFMILA-----TTSSTCAKLEEM 28
gi|PatB+ -----MATTKSFLVLFMILA-----TTSSTCATLGEM 28
gi|pentin1_phb -----MKSKMAMLLLLFCVLSNQLVAAFSTQAKASKDGNL 35
gi|5c9_phb -----MGSIGRGTANCATVPQPPSTGKL 24
gi|corn3_pep -----MGSIGRGTANCATVPQPPSTGKL 24
gi|corn2_pep -----MGSIGRGTANCATVPQPPSTGKL 24
gi|corn4_pep -----MGSIGRGTANCATVPQPPSTGKL 24
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gi|corn5_pep -----MGSIGRGTANCATVPQPPSTGKL 24
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gi|PatL+ VTVLSIDGGGIRGIIPATILEFLEGQLQEMDNNADARLADYFDVIGGTSTGGLLTAMITT 88
gi|PatA+ VTVLSIDGGGIRGIIPATILEFLEGQLQEMDNNADARLADYFDVIGGTSTGGLLTAMITT 88
gi|PatB+ VTVLSIDGGGIRGIIPATILEFLEGQLQEMDNNADARLADYFDVIGGTSTGGLLTAMITT 88
gi|pentin1_phb VTVLAIDGGGIRGIIPGVILKQLEATLQRWDSS--ARLAEYFDVAGTSTGGIITAILTA 93
gi|5c9_phb ITILSIDGGGIRGLIPATIIAYLEAKLQELDGP-DARIADYFDVIAGTSTGALLASMLAA 83
gi|corn3_pep ITILSIDGGGIRGLIPATIIAYLEAKLQELDGP-DARIADYFDVIAGTSTGALLASMLAA 83
gi|corn2_pep ITILSIDGGGIRGLIPATIIAYLEAKLQELDGP-DARIADYFDVIAGTSTGALLASMLAA 83
gi|corn4_pep ITILSIDGGGIRGLIPATIIAYLEAKLQELDGP-DARIADYFDVIAGTSTGALLASMLAA 83
gi|corn1_pep ITILSIDGGGIRGLIPATIIAYLEAKLQELDGP-DARIADYFDVIAGTSTGALLASMLAA 119
gi|corn5_pep ITILSIDGGGIRGLIPATIIAYLEAKLQELDGP-DARIADYFDVIAGTSTGALLASMLAA 83
::*:*****:*. *. *. *. *. *:*****:*****:*****:

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gi|PatFm PNENN--RPFAAAKEIVPFYFEHGPQIFNSR-----YWPIFWPKYDGKYLQVL 113
gi|PatIm PNENN--RPFAAAKDIVPFYFEHGPQIFNY-----SGSILGPMYDGKYLQVL 112
gi|PatL+ PNENN--RPFAAAKDIVPFYFEHGPQIFNY-----SGSILGPMYDGKYLQVL 134
gi|PatA+ PNENN--RPFAAAKDIVPFYFEHGPQIFNY-----SGSILGPMYDGKYLQVL 134
gi|PatB+ PNENN--RPFAAAKDIVPFYFEHGPQIFNS-----SGSIFGPMYDGKYFLQVL 134
gi|pentin1_phb PDPQNKDRPLYAAEEIIDFYIEHGSPSIFNKSTA-----CSLPGIFCPKYDGKYLQEI 146
gi|5c9_phb PDENN--RPLFAAKDLTTFYLENGPKIFPQKKAGLLTPLRNLLGLVRGPKYDGVFLHDKI 141
gi|corn3_pep PDENN--RPLFAAKDLTTFYLENGPKIFPQKKAGLLTPLRNLLGLVRGPKYDGVFLHDKI 141
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gi|corn1_pep PDENN--RPLFAAKDLTTFYLENGPKIFPQKKAGLLTPLRNLLGLVRGPKYDGVFLHDKI 177
gi|corn5_pep PDENN--RPLFAAKDLTTFYLENGPKIFPQKKAGLLTPLRNLLGLVRGPKYDGVFLHDKI 141
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Figure 9 (a)



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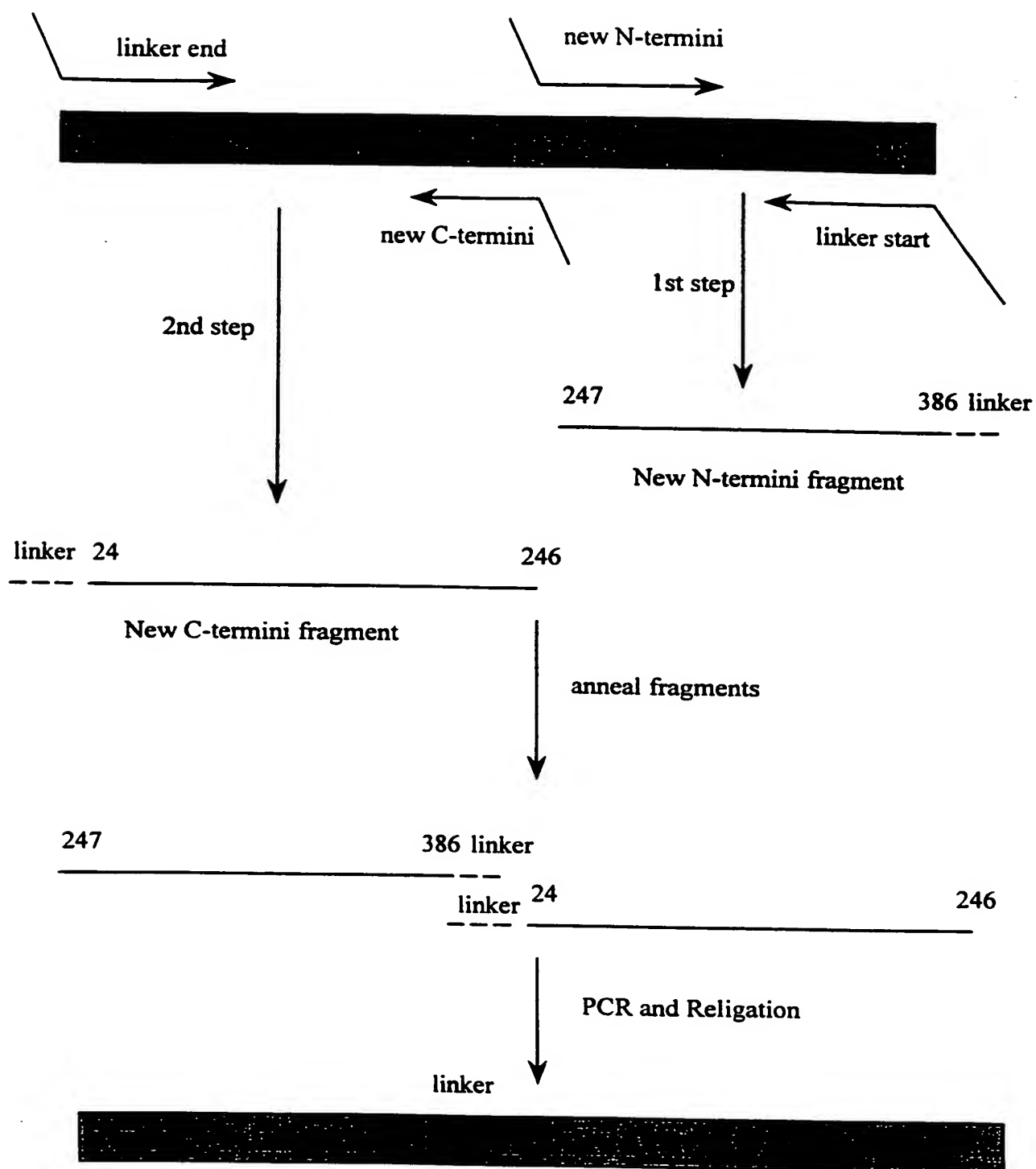
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gi|pentin1_phb      ALQPQNNYLRIQEYDLDP-----LESIDDASTENMENLEK 338
gi|5c9_phb          ALHCEKKYLRIQDDTLTGN-----ASSVDIATKENMESLIS 356
gi|corn3_pep        ALHCEKKYLRIQDDTLTGN-----ASSVDIATKENMESLIS 356
gi|corn2_pep        ALHCEKKYLRIQDDTLTGN-----ASSVDIATKENMESLIS 356
gi|corn4_pep        ALHCEKKYLRIQDDTLTGN-----ASSVDIATKENMESLIS 356
gi|corn1_pep        ALHCEKKYLRIQLYAGYFDWERIVRGHRHQGEHGVSDIDRPGAAQEASGESEHRAVR 416
gi|corn5_pep        ALHCEKKYLRIQLYAG-----337
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gi|patatin_mtc      VGENLLKKPVSEDNP-----ETYEEALKRFAKLLSDRKKLRANKASY----- 386
gi|PatFm            VGENLLKKPVSKDNP-----ETYEEALKRFAKLLSDRKKLRANKASY----- 365
gi|PatIm            VGETLLKKPVSRDSP-----ETYEEALKRFAKLLSDRKKLRANKASY----- 364
gi|PatL+            VGATLLKKPVSKDSP-----ETYEEALKRFAKLLSDRKKLRANKASY----- 386
gi|PatA+            VGETLLKKPVSKDSP-----ETYEEALKRFAKLLSDRKKLRANKASY----- 386
gi|PatB+            VGEKLLKKPVSKDSP-----ETYEEALKRFAKLLSDRKKLRANKASY----- 386
gi|pentin1_phb      VGQSLNNEPVKRMNLNT-FVVEETGEGTNAEALDRLAQIlyEEKITRGLGKISLEVDNID 397
gi|5c9_phb          IGQELLKKPVARVNIDTGLYESCDGEGTNAQSLADFAKQLSDERKLKSNLNSN----- 410
gi|corn3_pep        IGQELLKKPVARVNIDTGLYESCDGEGTNAQSLADFAKQLSDERKLKSNLNSN----- 410
gi|corn2_pep        IGQELLNKPVARVNIDTGLYESCEGEGTNAQSLADFAKQLSDERKLKSNLNSN----- 410
gi|corn4_pep        IGQELLNKPVARVNIDTGLYESCEGEGTNAQSLADFAKQLSDERKLKSNLNSN----- 410
gi|corn1_pep        VLRRGHKCTVASLRQATLRAQATQEQSQLQLINTSLSHSMCSFRFTVSFYFNFNSVCVL 476
gi|corn5_pep        -----

gi|patatin_mtc      -----
gi|PatFm            -----
gi|PatIm            -----
gi|PatL+            -----
gi|PatA+            -----
gi|PatB+            -----
gi|pentin1_phb      PYTERVRKLLF----- 408
gi|5c9_phb          -----
gi|corn3_pep        -----
gi|corn2_pep        -----
gi|corn4_pep        -----
gi|corn1_pep        CVLCVYQTFKFNQKKKKKKKKKKKKKKKRAA 508
gi|corn5_pep        -----

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**Figure 9 (c)**



**Figure 10**